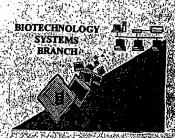


RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable

Application Serial Number: 09/509/65
Source: 1600 Pull Processed by STIC: 1/9/2002

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FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. War) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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- 2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/509,165

Input Set: A:\34810PCT.seq.txt
Output Set: N:\CRF3\01082002\1509165.raw

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4 <110> APPLICANT: ICOS Corporation, et al.
 6 <120> TITLE OF INVENTION: MACROPHAGE DERIVED CHEMOKINE (MDC), MDC ANALOGS, MDC
         INHIBITOR SUBSTANCES, AND USES THEREOF
 9 <130> FILE REFERENCE: 27866/34810PCT
11 <140> CURRENT APPLICATION NUMBER: US/09/509,165
12 <141> CURRENT FILING DATE: 1998-09-28
14 <150> PRIOR APPLICATION NUMBER: 09/067,447
15 <151> PRIOR FILING DATE: 1998-04-28
17 <150> PRIOR APPLICATION NUMBER: 08/939,107
18 <151> PRIOR FILING DATE: 1997-09-26
20 <150> PRIOR APPLICATION NUMBER: 08/660,542
                                                                Does Not Comply
21 <151> PRIOR FILING DATE: 1996-06-07
                                                            Corrected Diskette Needed
23 <150> PRIOR APPLICATION NUMBER: 08/558,658
24 <151> PRIOR FILING DATE: 1995-11-16
26 <150> PRIOR APPLICATION NUMBER: 08/479,620
27 <151> PRIOR FILING DATE: 1995-06-07
29 <160> NUMBER OF SEQ ID NOS: 46
31 <170> SOFTWARE: PatentIn Ver. 2.0
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ERRORED SEQUENCES

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397 <210> SEQ ID NO: 19
    398 <211> LENGTH: 99
                              sapiens
    399 <212> TYPE: PRT
    400 <213> ORGANISM: Homo (spapiens) - Hu MCP-1
A(2 401 <400> SEQUENCE: 19
    402 Met Lys Val Ser Ala Ala Leu Leu Cys Leu Leu Ile Ala Ala Thr
                    -20
                                        -15
    405 Phe Ile Pro Gln Gly Leu Ala Gln Pro Asp Ala Ile Asn Ala Pro Val
    408 Thr Cys Cys Tyr Asn Phe Thr Asn Arg Lys Ile Ser Val Gln Arg Leu
    409 10
                            15
                                                 20
    411 Ala Ser Tyr Arg Arg Ile Thr Ser Ser Lys Cys Pro Lys Glu Ala Val
    412
                        30
                                             35
    414 Ile Phe Lys Thr Ile Val Ala Lys Glu Ile Cys Ala Asp Pro Lys Gln
                                        50
    418 Lys Trp Val Gln Asp Ser Met Asp His Leu Asp Lys Gln Thr Gln Thr
    419
                60
                                    65
    421 Pro Lys Thr
    422
           75
   474 <210> SEQ ID NO: 22
    475 <211> LENGTH: 91
    476 <212> TYPE: PRT
    477 <213> ORGANISM: Homo sapiens - MIP-1 beta
    479 <220> FEATURE:
480 <400> SEQUENCE: 22
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(80) delete



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,165

DATE: 01/09/2002 TIME: 16:02:10

Input Set : A:\34810PCT.seq.txt

Output Set: N:\CRF3\01082002\I509165.raw

```
481 Met Lys Leu Cys Val Thr Val Leu Ser Leu Leu Met Leu Val Ala Ala
                    -20
                                        -15
     484 Phe Cys Ser Pro Ala Leu Ser Ala Pro Met Gly Ser Asp Pro Pro Thr
E--> 485 Liletel (-15)-5 (change to-5) 1 5
487 Ala Cys Cys Phe Ser Tyr Thr Arg Glu Ala Ser Ser Asn Phe Val Val
E--> 488 10
                                                20
     490 Asp Tyr Tyr Glu Thr Ser Ser Leu Cys Ser Ġln Pro Ala Val Val Phe
                                            35
     493 Gln Thr Lys Arg Ser Lys Gln Val Cys Ala Asp Pro Ser Glu Ser Trp
     496 Val Gln Glu Tyr Val Tyr Asp Leu Glu Leu Asn
                60
     500 <210> SEQ ID NO: 23
     501 <211> LENGTH: 92
     502 <212> TYPE: PRT
     503 <213> ORGANISM: Homo sapiens - MIP-1 alpha
     505 <220> FEATURE:
    506 <400> SEQUENCE: 23
     508 Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
                -20
                                    -15
     511 Leu Cys Asn Gln Phe Ser Ala Ser Leu Ala Ala Asp Thr Pro Thr Ala
     512 -5
     514 Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile Ala
                        15 43
     516 Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val Ile Phe
                    30
                                        35
     519 Leu Thr Lys Arg Ser Arg Gln Val Cys Ala Asp Pro Ser Glu Glu Trp
                45
                                    50
     522 Val Gln Lys Tyr Val Ser Asp Leu Glu Leu Ser Ala
                                65
            60
     526 <210> SEQ ID NO: 24
     527 <211> LENGTH: 96
     528 <212> TYPE: PRT
     529 <213> ORGANISM: Homo sapiens - I-309
    531 <220> FEATURE:
    532 <400> SEQUENCE: 24
     533 Met Gln Ile Ile Thr Thr Ala Leu Val Cys Leu Leu Ala Gly Met
                                   -15 <del>-15.</del>
     536 Trp Pro Glu Asp Val Asp Ser Lys Ser Met Gln Val Pro Phe Ser Arg
E-->537 -5 1 1
                                                5 5
     539 Cys Cys Phe Ser Phe Ala Glu Gln Glu Ile Pro Leu Arg Ala Ile Leu
                       15 15 20 -20
                                                                25 25-
    542 Cys Tyr Arg Asn Thr Ser Ser Ile Cys Ser Asn Glu Gly Leu Ile Phe
                    30 30 35 as
```

545 Lys Leu Lys Arg Gly Lys Glu Ala Cys Ala Leu Asp Thr Val Gly Trp

548 Val Gln Arg His Arg Lys Met Leu Arg His Cys Pro Ser Lys Arg Lys

65 65

50 -50

hunberry

40 40

55 .55-

70 70

su p.3

45 45

1114 <210> SEQ ID NO: 38

549 60 60

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,165

DATE: 01/09/2002

TIME: 16:02:10

Input Set : A:\34810PCT.seq.txt

Output Set: N:\CRF3\01082002\I509165.raw

1115 <211> LENGTH: 81 1116 <212> TYPE: PRT 1117 <213> ORGANISM: rat 1119 <400> SEQUENCE: 38 1120 Leu Val Leu Leu Ala Val Ala Leu Gln Thr Ser Asp Ala Gly Pro Tyr 1121 -10 -1 1123 Gly Ala Asn Val Glu Asp Ser Ile Cys Cys Gln Asp Tyr Ile Arg His 1124 1126 Pro Leu Pro Pro Arg Phe Val Lys Glu Phe Tyr Trp Thr Ser Lys Ser 1129 Cys Arg Lys Pro Gly Val Val Leu Ile Thr Ile Lys Asn Arg Asp Ile 1130 45 E--> 1132 Cys Ala Asp Pro (Xaa) Met Leu Trp Val Lys Lys Ile Leu His Lys Leu 1133

sel item 9 on Euro Sunnavy Sheet

Use of n and/or Xaa has been detected in the Sequence Listing.

Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

sel pp 4-7 for more enon

1/9/02

1135 Ala

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/509,165

DATE: 01/09/2002 TIME: 16:02:11

Input Set : A:\34810PCT.seq.txt
Output Set: N:\CRF3\01082002\1509165.raw

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L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:375 M:283 W: Missing Blank Line separator, <400> field identifier
L:401 M:283 W: Missing Blank Line separator, <400> field identifier
L:415 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:19
L:431 M:283 W: Missing Blank Line separator, <400> field identifier L:454 M:283 W: Missing Blank Line separator, <400> field identifier
L:480 M:283 W: Missing Blank Line separator, <400> field identifier
L:485 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:22
M:332 Repeated in SeqNo=22
L:506 M:283 W: Missing Blank Line separator, <400> field identifier
L:515 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:23
L:532 M:283 W: Missing Blank Line separator, <400> field identifier L:534 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:24
M:332 Repeated in SeqNo=24
L:603 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:603 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:603 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:606 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:606 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:606 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:609 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:609 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:609\ M:341\ W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:612 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:612 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
\text{L:612 M:341 W:} (46) "n" or "Xaa" used, for SEQ ID#:25
L:668 M:283 W: Missing Blank Line separator, <220> field identifier
L:670 M:283 W: Missing Blank Line separator, <400> field identifier
L:691 M:283 W: Missing Blank Line separator, <220> field identifier
L:715 M:283 W: Missing Blank Line separator, <220> field identifier
L:1082 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:37
L:1082 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1083\ M:258\ W: Mandatory Feature missing, <223> not found for SEQ ID#:37
L:1083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1105 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:37
L:1105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1107 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:37
L:1107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1109 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:37
\tt L\!:\!1109~M\!:\!341~W\!: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1111 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:37
L\!:\!1111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1132 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:38
L:1153 M:283 W: Missing Blank Line separator, <400> field identifier
L:1201 M:283 W: Missing Blank Line separator, <400> field identifier
L:1236 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:41
L:1240 M:283 W: Missing Blank Line separator, <220> field identifier
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/509,165

DATE: 01/09/2002 TIME: 16:02:11

Input Set : A:\34810PCT.seq.txt

Output Set: N:\CRF3\01082002\I509165.raw

L:1246 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:41 L:1246 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:41

L:1246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41